Please amend the application as follows:

In the claims:

Please cancel claims 118-159, 286, 288-290, 293, 294, 296, and 298-301 without prejudice or disclaimer.

Please replace claims 31, 47, 56, 63, 79, 95, 109, 291, 292, 302, 304, and 305 with the following amended claims:

- 31. (Amended) The nucleic acid molecule of claim 24 further comprising a nucleotide sequence heterologous to SEQ ID NO:1.
 - 47. (Thrice Amended) The nucleic acid molecule of claim 44 that further comprises a nucleotide sequence heterologous to SEQ ID NO:1.
 - 56. (Amended) An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:
 - (a) a nucleotide sequence encoding the full-length polypeptide encoded by the cDNA contained in clone HPHAE52 as deposited with the ATCC as accession number 97810;
 - (b) a nucleotide sequence encoding the full-length polypeptide, lacking the N-terminal methionine, which is encoded by the cDNA contained in clone HPHAE52 as deposited with the ATCC as accession number 97810;
 - (c) a nucleotide sequence encoding the mature polypeptide encoded by the cDNA contained in clone HPHAE52 as deposited with the ATCC as accession number 97810;
 - (d) a nucleotide sequence encoding the soluble extracellular domain encoded by the cDNA contained in clone HPHAE52 as deposited with the ATCC as accession number 97810, respectively; and
 - (e) a nucleotide sequence that is the complement of (a), (b), (c), or (d).
 - 63. (Amended) The nucleic acid molecule of claim 59 further comprising a nucleotide sequence heterologous to the cDNA contained in clone HPHAE52 as deposited with the ATCC as accession number 97810.

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79. (Amended) The nucleic acid molecule of claim 75 that further_comprises a nucleotide sequence heterologous to the cDNA contained in clone HPHAE52 as deposited with the ATCC as accession number 97810.



95. (Amended) The nucleic acid molecule of claim 88 further comprising a nucleotide sequence heterologous to SEQ ID NO:1.



109. (Amended) The nucleic acid molecule of claim 104 further comprising a nucleotide sequence heterologous to said cDNA clone.

291. (Amended) An isolated polynucleotide comprising a nucleotide sequence that is at least 95% identical to a nucleotide sequence encoding amino acid residues 31-300 of SEQ ID NO:2 wherein said polynucleotide encodes a polypeptide that binds Fas ligand.



292. (Amended) An isolated polynucleotide comprising a nucleotide sequence that is at least 95% identical-to a nucleotide sequence encoding amino acid residues 31-283 of SEQ ID NO: 2 wherein said polynucleotide encodes a polypeptide that binds Fas ligand.



302. (Amended) An expression vector for the production of a polypeptide comprising amino acids 31-300 of SEQ ID NO:2 comprising a polynucleotide that encodes amino acids 31-300 of SEQ ID NO:2 operably associated with a regulatory element that controls expression of said polynucleotide.



304. (Amended) A method of for producing a polypeptide comprising amino acids 31-300 of SEQ ID NO:2, comprising culturing the host cell of claim 303 under conditions sufficient for the production of said polypeptide and recovering said polypeptide from the culture.

305. (Amended) A process for producing a cell which produces a polypeptide comprising amino acids 31-300 of SEQ ID NO:2, comprising transforming or transfecting a host cell with the expression vector of claim 302 such that the host cell, under appropriate culture conditions, produces said polypeptide.

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